

Additional file 7:

Figure S7. Hotspot densities are reduced in gene clusters with silenced expression in spermatocytes. A representative region on mouse Chromosome 3 is shown. The first and second row represent domains enriched (above the line) or depleted (below the line) for H3K9me3 and H3K9me2, respectively; third row – positions of all Affinity-seq sites; fourth row – positions of Affinity-seq sites detected at hotspots *in vivo*; fifth row – positions of genes not expressed in spermatocytes.

